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Phe Leu Gly Cys His Met Val Ala Ile Val Pro Asp Leu Leu Gln Gly 65 70 75 80

Arg Leu Asp Phe Pro Gly Phe Val Gln Thr Ser Leu Ala Thr Leu Arg 85 90 95

Phe Phe Cys Tyr Ile Val Gly Leu Ser Leu Leu Ala Ala Val Ser Val

Glu Gln Cys Leu Ala Ala Leu Phe Pro Ala Trp Tyr Ser Cys Arg Arg 115 120 125

Pro Arg His Leu Thr Thr Cys Val Cys Ala Leu Thr Trp Ala Leu Cys 130 135 140

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His Asn Leu Val Ala Ser Thr Ala Val Ser Asp Glu Leu Val Ala Ala 50 55 60

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Leu Pro Ala Val Gly Trp His Asp Thr Ser Glu Arg Phe Tyr Thr His 100 105 110

Gly Cys Arg Phe Ile Val Ala Glu Ile Gly Leu Gly Phe Gly Val Cys 115 120 125

Phe Leu Leu Val Gly Gly Ser Val Ala Met Gly Val Ile Cys Thr 130 135 140

Ala Ile Ala Leu Phe Gln Thr Leu Ala Val Gln Val Gly Arg Gln Ala 145 150 155 160

Asp Arg Arg Ala Phe Thr Val Pro Thr Ile Val Val Glu Asp Ala Gln 165 170 175

Gly Lys Arg Arg Ser Ser Ile Asp Gly Ser Glu Pro Ala Lys Thr Ser 180 185 190

Leu Gln Thr Thr Gly Leu Val Thr Thr Ile Val Phe Ile Tyr Asp Cys 195 200 205

Leu Met Gly Phe Pro Val Leu Val 210 215

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<210> 23
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<213> Homo sapiens

<400> 23

Met Ser Asp Glu Arg Arg Leu Pro Gly Ser Ala Val Gly Trp Leu Val

Cys Gly Gly Leu Ser Leu Leu Ala Asn Ala Trp Gly Ile Leu Ser Val 20 25 30

Gly Ala Lys Gln Lys Lys Trp Lys Pro Leu Glu Phe Leu Leu Cys Thr 35 40 45

Leu Ala Ala Thr His Met Leu Asn Val Ala Val Pro Ile Ala Thr Tyr 50 55 60

Ser Val Val Gln Leu Arg Arg Gln Arg Pro Asp Phe Glu Trp Asn Glu 65 70 75 80

Gly Leu Cys Lys Val Phe Val Ser Thr Phe Tyr Thr Leu Thr Leu Ala 85 90 95

Thr Cys Phe Ser Val Thr Ser Leu Ser Tyr His Arg Met Trp Met Val 100 105 110

Cys Trp Pro Val Asn Tyr Arg 115

<210> 24

<211> 330

<212> PRT

<213> Homo sapiens

<400> 24

Met Asp Pro Thr Thr Pro Ala Trp Gly Thr Glu Ser Thr Thr Val Asn 1 5 10 15

Gly Asn Asp Gln Ala Leu Leu Leu Cys Gly Lys Glu Thr Leu Ile 20 25 30

Pro Val Phe Leu Ile Leu Phe Ile Ala Leu Val Gly Leu Val Gly Asn 35 40 45

Gly Phe Val Leu Trp Leu Leu Gly Phe Arg Met Arg Arg Asn Ala Phe 50 55 60

Ser Val Tyr Val Leu Ser Leu Ala Gly Ala Asp Phe Leu Phe Leu Cys 65 70 75 80

Phe Gln Ile Ile Asn Cys Leu Val Tyr Leu Ser Asn Phe Phe Cys Ser 85 90 95

Ile Ser Ile Asn Phe Pro Ser Phe Phe Thr Thr Val Met Thr Cys Ala 100 105 110

Tyr Leu Ala Gly Leu Ser Met Leu Ser Thr Val Ser Thr Glu Arg Cys

<211> 119

<212> PRT

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Leu	Ser 130	Val	Leu	Trp		Ile 135	Trp	Tyr	Arg	Cys	Arg 140	Arg	Pro	Arg	His
Leu	Ser	Ala	Val	Val	Cys	Val	Leu	Leu	Trp	Ala	Leu	Ser	Leu	Leu	Leu

Ser Ile Leu Glu Gly Lys Phe Cys Gly Phe Leu Phe Ser Asp Gly Asp

Ser Gly Trp Cys Gln Thr Phe Asp Phe Ile Thr Ala Ala Trp Leu Ile 180 185 190

Phe Leu Phe Met Val Leu Cys Gly Ser Ser Leu Ala Leu Leu Val Arg 195 200 205

Ile Leu Cys Gly Ser Arg Gly Leu Pro Leu Thr Arg Leu Tyr Leu Thr 210 215 220

Ile Leu Leu Thr Val Leu Val Phe Leu Leu Cys Gly Leu Pro Phe Gly 225 230 235 240

Ile Gln Trp Phe Leu Ile Leu Trp Ile Trp Lys Asp Ser Asp Val Leu 245 250 255

Phe Cys His Ile His Pro Val Ser Val Val Leu Ser Ser Leu Asn Ser 260 265 270

Ser Ala Asn Pro Ile Ile Tyr Phe Phe Val Gly Ser Phe Arg Lys Gln 275 280 285

Trp Arg Leu Gln Gln Pro Ile Leu Lys Leu Ala Leu Gln Arg Ala Leu 290 295 300

Gln Asp Ile Ala Glu Val Asp His Ser Glu Gly Cys Phe Arg Gln Gly 305 310 315 320

Thr Pro Glu Met Ser Arg Ser Ser Leu Val 325 330

<210> 25

<211> 371

<212> PRT

<213> Homo sapiens

<400> 25

Met Pro Ala Asn Phe Thr Glu Gly Ser Phe Asp Ser Ser Gly Thr Gly 1 5 10 15

Gln Thr Leu Asp Ser Ser Pro Val Ala Cys Thr Glu Thr Val Thr Phe 20 25 30

Thr Glu Val Val Glu Gly Lys Glu Trp Gly Ser Phe Tyr Tyr Ser Phe 35 40 45

Lys Thr Glu Gln Leu Ile Thr Leu Trp Val Leu Phe Val Phe Thr Ile 50 55 60

Val Gly Asn Ser Val Val Leu Phe Ser Thr Trp Arg Arg Lys Lys

Ser Arg Met Thr Phe Phe Val Thr Gln Leu Ala Ile Thr Asp Ser Phe 85 Thr Gly Leu Val Asn Ile Leu Thr Asp Ile Ile Trp Arg Phe Thr Gly 100 105 Asp Phe Thr Ala Pro Asp Leu Val Cys Arg Val Val Arg Tyr Leu Gln 120 Val Val Leu Leu Tyr Ala Ser Thr Tyr Val Leu Val Ser Leu Ser Ile Asp Arg Tyr His Ala Ile Val Tyr Pro Met Lys Phe Leu Gln Gly Glu 150 Lys Gln Ala Arg Val Leu Ile Val Ile Ala Trp Ser Leu Ser Phe Leu Phe Ser Ile Pro Thr Leu Ile Ile Phe Gly Lys Arg Thr Leu Ser Asn Gly Glu Val Gln Cys Trp Ala Leu Trp Pro Gly Asp Ser Tyr Trp Thr Pro Tyr Met Thr Ile Val Ala Phe Leu Val Tyr Phe Ile Pro Leu Thr Ile Ile Ser Ile Met Tyr Gly Ile Val Ile Arg Thr Ile Trp Ile Lys 235 Ser Lys Thr Tyr Glu Thr Val Ile Ser Asn Cys Ser Asp Gly Lys Leu Cys Ser Ser Tyr Asn Arg Gly Leu Ile Ser Lys Ala Lys Ile Lys Ala 265 Ile Lys Tyr Ser Ile Ile Ile Leu Ala Phe Ile Cys Cys Trp Ser Pro Tyr Phe Leu Phe Asp Ile Leu Asp Asn Phe Asn Leu Leu Pro Asp Thr Glm Glu Arg Phe Tyr Ala Ser Val Ile Ile Gln Asn Leu Pro Ala 310 Leu Asn Ser Ala Ile Asn Pro Pro Ile Tyr Cys Val Phe Ser Ser Ser 325 Ile Ser Phe Pro Cys Arg Glu Gln Arg Ser Gln Asp Ser Arg Met Thr 345 340 Phe Arg Glu Arg Thr Glu Arg His Glu Met Gln Ile Leu Ser Lys Pro 360 Glu Phe Ile 370

<210> 26 <211> ·393

<212> PRT

<213> Homo sapiens

<400> 26

Met Glu Thr Thr Met Gly Phe Met Asp Asp Asn Ala Thr Asn Thr Ser 1 10 15

Thr Ser Phe Leu Ser Val Leu Asn Pro His Gly Ala His Ala Thr Ser 20 25 30

Phe Pro Phe Asn Phe Ser Tyr Ser Asp Tyr Asp Met Pro Leu Asp Glu 35 40 45

Asp Glu Asp Val Thr Asn Ser Arg Thr Phe Phe Ala Ala Lys Ile Val 50 55 60

Ile Gly Met Ala Leu Val Gly Ile Met Leu Val Cys Gly Ile Gly Asn 65 70 75 80

Phe Ile Phe Ile Ala Ala Leu Val Arg Tyr Lys Lys Leu Arg Asn Leu 85 90 95

Thr Asn Leu Leu Ile Ala Asn Leu Ala Ile Ser Asp Phe Leu Val Ala 100 105 110

Ile Val Cys Cys Pro Phe Glu Met Asp Tyr Tyr Val Val Arg Gln Leu 115 120 125

Ser Trp Glu His Gly His Val Leu Cys Thr Ser Val Asn Tyr Leu Arg 130 135 140

Thr Val Ser Leu Tyr Val Ser Thr Asn Ala Leu Leu Ala Ile Ala Ile 145 150 155 160

Asp Arg Tyr Leu Ala Ile Val His Pro Leu Arg Pro Arg Met Lys Cys 165 170 175

Gln Thr Ala Thr Gly Leu Ile Ala Leu Val Trp Thr Val Ser Ile Leu 180 185 190

Ile Ala Ile Pro Ser Ala Tyr Phe Thr Thr Glu Thr Val Leu Val Ile 195 200 205

Val Lys Ser Gln Glu Lys Ile Phe Cys Gly Gln Ile Trp Pro Val Asp 210 215 220

Gln Gln Leu Tyr Tyr Lys Ser Tyr Phe Leu Phe Ile Phe Gly Ile Glu 225 230 235 240

Phe Val Gly Pro Val Val Thr Met Thr Leu Cys Tyr Ala Arg Ile Ser 245 250 255

Arg Glu Leu Trp Phe Lys Ala Val Pro Gly Phe Gln Thr Glu Gln Ile 260 265 270

Arg Lys Arg Leu Arg Cys Arg Arg Lys Thr Val Leu Val Leu Met Cys 275 280 285

Ile Leu Thr Ala Tyr Val Leu Cys Trp Ala Pro Phe Tyr Gly Phe Thr 290 295 300

Ile Val Arg Asp Phe Phe Pro Thr Val Phe Val Lys Glu Lys His Tyr 305 310 315 320

Leu Thr Ala Phe Tyr Ile Val Glu Cys Ile Ala Met Ser Asn Ser Met 325 330 335

Ile Asn Thr Leu Cys Phe Val Thr Val Lys Asn Asp Thr Val Lys Tyr 340 345 350

Phe Lys Lys Ile Met Leu Leu His Trp Lys Ala Ser Tyr Asn Gly Gly 355 360 365

Lys Ser Ser Ala Asp Leu Asp Leu Lys Thr Ile Gly Met Pro Ala Thr 370 375 380

Glu Glu Val Asp Cys Ile Arg Leu Lys 385 390

<210> 27

<211> 389

<212> PRT

<213> Homo sapiens

<400> 27

Met Gly Phe Met Asp Asp Asn Ala Thr Asn Thr Ser Thr Ser Phe Leu

5 10 15

Ser Val Leu Asn Pro His Gly Ala His Ala Thr Ser Phe Pro Phe Asn 20 25 30

Phe Ser Tyr Ser Asp Tyr Asp Met Pro Leu Asp Glu Asp Glu Asp Val 35 40 45

Thr Asn Ser Arg Thr Phe Phe Ala Ala Lys Ile Val Ile Gly Met Ala 50 55 60

Leu Val Gly Ile Met Leu Val Cys Gly Ile Gly Asn Phe Ile Phe Ile 65 70 75 80

Ala Ala Leu Val Arg Tyr Lys Lys Leu Arg Asn Leu Thr Asn Leu Leu 85 90 95

Ile Ala Asn Leu Ala Ile Ser Asp Phe Leu Val Ala Ile Val Cys Cys
100 105 110

Pro Phe Glu Met Asp Tyr Tyr Val Val Arg Gln Leu Ser Trp Glu His 115 120 125

Gly His Val Leu Cys Thr Ser Val Asn Tyr Leu Arg Thr Val Ser Leu 130 135 140

Tyr Val Ser Thr Asn Ala Leu Leu Ala Ile Ala Ile Asp Arg Tyr Leu 145 150 155 160

Ala Ile Val His Pro Leu Arg Pro Arg Met Lys Cys Gln Thr Ala Thr 165 170 175

Gly Leu Ile Ala Leu Val Trp Thr Val Ser Ile Leu Ile Ala Ile Pro 180 185 190

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Ser Ala Tyr Phe Thr Thr Glu Thr Val Leu Val Ile Val Lys Ser Gln 195 205 Glu Lys Ile Phe Cys Gly Gln Ile Trp Pro Val Asp Gln Gln Leu Tyr
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210 215 220

Tyr Lys Ser Tyr Phe Leu Phe Ile Phe Gly Ile Glu Phe Val Gly Pro

Val Val Thr Met Thr Leu Cys Tyr Ala Arg Ile Ser Arg Glu Leu Trp

230

Phe Lys Ala Val Pro Gly Phe Gln Thr Glu Gln Ile Arg Lys Arg Leu 260 265 270

Arg Cys Arg Arg Lys Thr Val Leu Val Leu Met Cys Ile Leu Thr Ala 275 280 285

Tyr Val Leu Cys Trp Ala Pro Phe Tyr Gly Phe Thr Ile Val Arg Asp 290 295 300

Phe Phe Pro Thr Val Phe Val Lys Glu Lys His Tyr Leu Thr Ala Phe 305 310 315 320

Tyr Ile Val Glu Cys Ile Ala Met Ser Asn Ser Met Ile Asn Thr Leu 325 330 335

Cys Phe Val Thr Val Lys Asn Asp Thr Val Lys Tyr Phe Lys Lys Ile 340 345 350

Met Leu Leu His Trp Lys Ala Ser Tyr Asn Gly Gly Lys Ser Ser Ala 355 360 365

Asp Leu Asp Leu Lys Thr Ile Gly Met Pro Ala Thr Glu Glu Val Asp 370 375 380

Cys Ile Arg Leu Lys 385

<210> 28

<211> 9

<212> PRT

<213> Synthetic Substrate

<400> 28

Ala Pro Arg Thr Pro Gly Gly Arg Arg

<210> 29

<211> 27

<212> DNA

<213> Artificial

<220>

<223> Novel Sequence

<400> 29

ttcaaagctt atggatccaa ccacccc

27

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<220> <223>	Novel Sequence	
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	31 19 DNA Artificial	
<220> <223>	Novel Sequence	
<400> taggca	31 cagg tcatcacag	19
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<400> ttggac	32 egcca ggaaggtg	18
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<220> <223>	Novel Sequence	
<400> gcctgg	33 gagee tgtetttet gttete	26
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<400> gtaga	34 tgagg gggttgatgg cactattc	28

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<220> <223>	Novel Sequence	
<400> gatag	36 ccttg atttttgcct ttgagatg	28
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<400> gcagc	37 agaca atggccacca ggaagtcaga	30
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<211> <212> <213>	19 DNA Artificial					
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tctcaga	accc tctccccaca	tctcctggtc	cctgccccca	cctggcgtac	agggaccagc	180
cccacg	gaag gctcttgagg	ccaggtaacc	atggggaggg	gaggaatggg	gacaccttcc	240
tcctgag	gtgt cttagggaag	agaagcttag	gtcaggtggc	tgagggtgga	aatgagagag	300
gggtct	cete etggagggte	tcaccattcc	cttggtcacc	cacccaactc	tcatctcccc	360
tgatgt	gggg aggagcaggg	ggcatggatt	cctgagcccc	agactcaact	gttgtggttt	420
acaggg	gcat caggagagag	agcgagcaga	acacactcct	gcagcatccc	ctggcccccc	480
gccccat	tgat ggagcccaga	gaagctggac	agcacgtggg	ggccgccaac	ggcgcccagg	540
aggatg	tggc cttcaacctc	atcatcctgt	ccctcaccga	ggggctcggc	ctcggtgggc	600
tgctgg	ggaa tggggcagtc	ctctggctgc	tcagctccaa	tgtctacaga	aaccccttcg	660
ccatcta	acct cctggacgtg	gcctgcgcgg	atctcatctt	ccttggctgc	cacatggtgg	720
ccatcg	tccc cgacttgctg	caaggccggc	tggacttccc	gggcttcgtg	cagaccagcc	780
tggcaa	cgct gcgcttcttc	tgctacatcg	tgggcctgag	tctcctggcg	gccgtcagcg	840
tggagc	agtg cctggccgcc	ctcttcccag	cctggtactc	gtgccgccgc	ccacgccacc	900
tgacca	cctg tgtgtgcgcc	ctcacctggg	ccctctgcct	gctgctgcac	ctgctgctca	960
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tggtgg	cage ggtgctgctg	gctctgctgt	gttgcaccat	gtgtggggcc	agccttatgc	1080
tgctgc	tgcg ggtggagcga	ggcccccagc	ggcccccacc	ccggggcttc	cctgggctca	1140
tcctcc	tcac cgtcctcctc	ttcctcttct	gcggcctgcc.	cttcggcatc	tactggctgt	1200
cccgga	ácct gctctggtac	atcccccact	acttctacca	cttcagcttc	ctcatggccg	1260
aaataa	anta cacaaccasa	cccatcatct	acttctqcct	aaacsatacc	cadddccdca	1320

ggctgcccct	ccggctggtc	ctccagcgag	cgctgggaga	cgaggctgag	cradadacca	1380
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ccccagctgc	agcccccgtg	aggcaagagg	gtgacgtggg	gaaggtggtg	gggtcagagg	1500
ctggggccag	ccggacctgg	aggaggcctt	ggtgggtgac	ccggtcatgt	gctgtcaaag	1560
ttgtgaccct	tggtctggag	catgaggctc	ccctgggagg	cagctggaaa	gg	1612

<210> 42

<211> 530

<212> PRT

<213> Homo sapiens

<400> 42

Val Ser Arg Asp Gly Ala Ile Ala Leu Pro Gly Ala Thr Glu Pro Asp 1 5 10 15

Ser Ile Ser Lys Lys Lys Arg Pro Phe Gly Ser Arg His His Gln Gln 20 25 30

Gly Ala Pro Trp Val Ser Asp Pro Leu Pro Thr Ser Pro Gly Pro Cys
35 40 45

Pro His Leu Ala Tyr Arg Asp Gln Pro His Gly Arg Leu Leu Arg Pro 50 55 60

Gly Asn His Gly Glu Gly Arg Asn Gly Asp Thr Phe Leu Leu Ser Val 65 70 75 80

Leu Gly Lys Arg Ser Leu Gly Gln Val Ala Glu Gly Gly Asn Glu Arg 85 90 95

Gly Val Ser Ser Trp Arg Val Ser Pro Phe Pro Trp Ser Pro Thr Gln
100 105 110

Leu Ser Ser Pro Leu Met Trp Gly Gly Ala Gly Gly Met Asp Ser Ala 115 120 125

Pro Asp Ser Thr Val Val Val Tyr Arg Gly Ile Arg Arg Glu Ser Glu 130 135 140

Gln Asn Thr Leu Leu Gln His Pro Leu Ala Pro Arg Pro Met Met Glu 145 150 155 160

Pro Arg Glu Ala Gly Gln His Val Gly Ala Ala Asn Gly Ala Gln Glu 165 170 175

Asp Val Ala Phe Asn Leu Ile Ile Leu Ser Leu Thr Glu Gly Leu Gly 180 185 190

Leu Gly Gly Leu Leu Gly Asn Gly Ala Val Leu Trp Leu Leu Ser Ser 195 200 205

Asn Val Tyr Arg Asn Pro Phe Ala Ile Tyr Leu Leu Asp Val Ala Cys 210 215 220

Ala Asp Leu Ile Phe Leu Gly Cys His Met Val Ala Ile Val Pro Asp





Leu Leu Gln Gly Arg Leu Asp Phe Pro Gly Phe Val Gln Thr Ser Leu 245 250 255

Ala Thr Leu Arg Phe Cys Tyr Ile Val Gly Leu Ser Leu Leu Ala Ala 260 265 270

Val Ser Val Glu Gln Cys Leu Ala Ala Leu Phe Pro Ala Trp Tyr Ser 275 280 285

Cys Arg Arg Pro Arg His Leu Thr Thr Cys Val Cys Ala Leu Thr Trp 290 295 300

Ala Leu Cys Leu Leu Leu His Leu Leu Leu Ser Gly Ala Cys Thr Gln 305 310 315

Phe Phe Gly Glu Pro Ser Arg His Leu Cys Arg Thr Leu Trp Leu Val 325 330 335

Ala Ala Val Leu Leu Ala Leu Leu Cys Cys Thr Met Cys Gly Ala Ser 340 345 350

Leu Met Leu Leu Arg Val Glu Arg Gly Pro Gln Arg Pro Pro Pro 355 360 365

Arg Gly Phe Pro Gly Leu Ile Leu Leu Thr Val Leu Leu Phe Leu Phe. 370 375 380

Cys Gly Leu Pro Phe Gly Ile Tyr Trp Leu Ser Arg Asn Leu Leu Trp 385 390 395

Tyr Ile Pro His Tyr Phe Tyr His Phe Ser Phe Leu Met Ala Ala Val

His Cys Ala Ala Lys Pro Val Val Tyr Phe Cys Leu Gly Ser Ala Gln 420 425 430

Gly Arg Arg Leu Pro Leu Arg Leu Val Leu Gln Arg Ala Leu Gly Asp 435 440 445

Glu Ala Glu Leu Gly Ala Val Arg Glu Thr Ser Arg Arg Gly Leu Val 450 455 460

Asp Ile Ala Ala Ala Leu Gly Pro Pro Thr Pro Ala Ala Ala Pro Val

Arg Gln Glu Gly Asp Val Gly Lys Val Val Gly Ser Glu Ala Gly Ala

Ser Arg Thr Trp Arg Arg Pro Trp Trp Val Thr Arg Ser Cys Ala Val 500 505 510

Lys Val Val Thr Leu Gly Leu Glu His Glu Ala Pro Leu Gly Gly Ser 515 520 525

Trp Lys 530

<210> 43 <211> 1612 <212> DNA <213> Homo sapiens

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<210> 44 <211> 311

<212> PRT

<213> Homo sapiens

<400> 44

Met Met Glu Pro Arg Glu Ala Gly Gln His Val Gly Ala Ala Asn Gly
1 5 10 15

Ala Gln Glu Asp Val Ala Phe Asn Leu Ile Ile Leu Ser Leu Thr Glu 20 · 25 30

Gly Leu Gly Leu Gly Gly Leu Leu Gly Asn Gly Ala Val Leu Trp Leu 35 40 45

Leu Ser Ser Asn Val Tyr Arg Asn Pro Phe Ala Ile Tyr Leu Leu Asp 50 55 60

Val Ala Cys Ala Asp Leu Ile Phe Leu Gly Cys His Met Val Ala Ile 65 70 75 80

Val Pro Asp Leu Leu Gln Gly Arg Leu Asp Phe Pro Gly Phe Val Gln 85 90 95

Thr Ser Leu Ala Thr Leu Arg Phe Cys Tyr Ile Val Gly Leu Ser Leu 100 105 110

Leu Ala Ala Val Ser Val Glu Gln Cys Leu Ala Ala Leu Phe Pro Ala 115 120 125

Trp Tyr Ser Cys Arg Arg Pro Arg His Leu Thr Thr Cys Val Cys Ala

Leu Thr Trp Ala Leu Cys Leu Leu Leu His Leu Leu Leu Ser Gly Ala 145 150 150 160

Cys Thr Gln Phe Phe Gly Glu Pro Ser Arg His Leu Cys Arg Thr Leu 165 170 175

Trp Leu Val Ala Ala Val Leu Leu Ala Leu Leu Cys Cys Thr Met Cys
180 185 190

Gly Ala Ser Leu Met Leu Leu Leu Arg Val Glu Arg Gly Pro Gln Arg 195 200 205

Pro Pro Pro Arg Gly Phe Pro Gly Leu Ile Leu Leu Thr Val Leu Leu 210 215 220

Phe Leu Phe Cys Gly Leu Pro Phe Gly Ile Tyr Trp Leu Ser Arg Asn 225 230 235 240

Leu Leu Trp Tyr Ile Pro His Tyr Phe Tyr His Phe Ser Phe Leu Met 245 250 255

Ala Ala Val His Cys Ala Ala Lys Pro Val Val Tyr Phe Cys Leu Gly 260 265 270

Ser Ala Gln Gly Arg Arg Leu Pro Leu Arg Leu Val Leu Gln Arg Ala 275 280 285

Leu Gly Asp Glu Ala Glu Leu Gly Ala Val Arg Glu Thr Ser Arg Arg 290 295 300



Gly Leu Val Asp Ile Ala Ala 305 310

<210> 45

<211> 939

<212> DNA

<213> Homo sapiens

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<210> 46

<211> 311

<212> PRT

<213> Homo sapiens

<400> 46

Met Met Glu Pro Arg Glu Ala Gly Gln His Val Gly Ala Ala Asn Ser 1 5 10 15

Ala Gln Glu Asp Val Ala Phe Asn Leu Ile Ile Leu Ser Leu Thr Glu 20 25 30

Gly Leu Gly Leu Gly Leu Leu Gly Asn Gly Ala Val Leu Trp Leu 35 40 45

Leu Ser Ser Asn Val Tyr Arg Asn Pro Phe Ala Ile Tyr Leu Leu Asp 50 55 60



Val Ala Cys Ala Asp Leu Ile Phe Leu Gly Cys His Met Val Ala Ile 65 70 75 80

Val Pro Asp Leu Leu Gln Gly Arg Leu Asp Phe Pro Gly Phe Val Gln 85 90 95

Thr Ser Leu Ala Thr Leu Arg Phe Cys Tyr Ile Val Gly Leu Ser Leu 100 105 110

Leu Ala Ala Val Ser Val Glu Gln Cys Leu Ala Ala Leu Phe Pro Ala 115 120 125

Trp Tyr Ser Cys Arg Arg Pro Arg His Leu Thr Thr Cys Val Cys Ala 130 135 140

Leu Thr Trp Ala Leu Cys Leu Leu Leu His Leu Leu Leu Ser Gly Ala 145 150 155 160

Cys Thr Gln Phe Phe Gly Glu Pro Ser Arg His Leu Cys Arg Thr Leu 165 170 175

Trp Leu Val Ala Ala Val Leu Leu Ala Leu Leu Cys Cys Thr Met Cys 180 185 190

Gly Ala Ser Leu Met Leu Leu Leu Arg Val Glu Arg Gly Pro Gln Arg 195 200 205

Pro Pro Pro Arg Gly Phe Pro Gly Leu Ile Leu Leu Thr Val Leu Leu 210 215 220

Phe Leu Phe Cys Gly Leu Pro Phe Gly Ile Tyr Trp Leu Ser Arg Asn 225 230 235

Leu Leu Trp Tyr Ile Pro His Tyr Phe Tyr His Phe Ser Phe Leu Met 245 250 255

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